

GeDiPNet Cluster 276 -- Pairs Within Cluster

Similarity: Jaccard index on shared curated genes (disease_gdp). Significance: upper-tail hypergeometric test, Benjamini-Hochberg FDR-corrected across all tested pairs. See scripts/build_related.php on gedipnet.bicnirrh.res.in. Data as of: 2026-10-07 03:00:13. · Rows: 9 · Generated: 2026-10-07 03:11:09

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Congenital cataract hypertrophic cardiomyopathy mitochondrial myopathy syndrome	Sengers syndrome	0.60000	3	6.575e-12	7.736e-11
Congenital cataract hypertrophic cardiomyopathy mitochondrial myopathy syndrome	mitochondrial DNA depletion syndrome 12B (cardiomyopathic type), autosomal recessive	0.25000	1	1.948e-4	5.278e-4
Congenital cataract hypertrophic cardiomyopathy mitochondrial myopathy syndrome	Triokinase and fmn cyclase deficiency	0.25000	1	1.948e-4	5.278e-4
mitochondrial DNA depletion syndrome 12B (cardiomyopathic type), autosomal recessive	Sengers syndrome	0.20000	1	2.598e-4	6.403e-4
Sengers syndrome	Triokinase and fmn cyclase deficiency	0.20000	1	2.598e-4	6.403e-4
Trichohepatoenteric syndrome	trichohepatoenteric syndrome 2	0.20000	1	2.598e-4	6.403e-4
Trichohepatoenteric syndrome	trichohepatoenteric syndrome 1	0.20000	1	2.598e-4	6.403e-4
Congenital cataract hypertrophic cardiomyopathy mitochondrial myopathy syndrome	Trichohepatoenteric syndrome	0.14286	1	7.792e-4	1.386e-3
Sengers syndrome	Trichohepatoenteric syndrome	0.12500	1	1.039e-3	1.720e-3